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SPECIATION IN THE AGATHYMUS (Megathymidae)

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I CAN REMEMBER BACK in the past when we used to collect butterflies with a net and killing jar and then there were fifteen recognized species in two genera in the family Megathymidae. Today we have 47 species in five genera in that family and the net has long since been retired after we discovered that it was much more efficient to collect these "critters" with a fox hole pick, drain spade or sharp knife. Once when we had available a few hundred specimens of these insects the taxonomy seemed comparatively simple, and now that we have several thousand specimens available for study complications present themselves.

In making a study of speciation in the Agathymus we must recognize a number of species complexes that exist in that genus. They are the *neumoegeni*, *chisosensis*, *hoffmanni*, *evansi*, *aryxna*, *baueri*, *alliae*, *stephensi*, *polingi*, *mariae*, *remingtoni*, *rethon*, and *indecisa* complexes. Of these the following are very complicated and indicate a considerable amount of evolution is in progress: *neumoegeni*, *hoffmanni*, *aryxna*, *mariae*, and *remingtoni*, and it is with this group that I would like to go into with some detail.

Agathymus speciation particularly in the mariae complex has intrigued me for a number of reasons, one of which has been the method by which new species apparently developed during the past ages from a mariae-like prototype. In making a detailed study of this species complex in the United States a number of very interesting observations were encountered some of which I would like to discuss briefly. One of the first things that was noticed about mariae was its close association with its larval food plant Agave Lecheguilla Torr. During the number of

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years that I have collected mariae and other related species I have never found a specimen in any other species of Agave other than members of the lecheguilla complex, with the possible exception of a Mexican species that feeds in Agave falcata Engelm. Indications are that at one time *lecheguilla* occurred extensively from the Edwards Plateau in Texas over into New Mexico and well down into Mexico. Fossil seeds have been found in areas of Texas now completely devoid of these plants. In this vast expanse a prototype of our modern species was well established with a gene pool common to all areas. Conditions arose resulting in the eradication of *lecheguilla* plants in many sections of this great area thus disrupting the gene pool. During the many years that followed mutations occurred in the restricted populations resulting in the production of individuals that differed from the original species. These isolated populations gradually changed both biologically and morphologically into five separate species in the Texas area. Mariae was the first described in this complex and it was found at El Paso, in the Franklin Mountains, and for many years was considered to be the only species present all over southwestern Texas. After carefully studying specimens from 32 locations in Texas and New Mexico it was found that there was actually five different species going under the name of mariae. Chinatiensis Freeman, in the Chinati Mountains and near vicinity; lajitaensis Freeman, in the Lajita region of the Big Bend; rindgei Freeman, and gilberti Freeman, in the general vicinity of Del Rio; and mariae (B. & B.), from the Big Bend and McCamev regions westward and north to Carlsbad. New Mexico and El Paso, Texas. In no area do more than two of these species occur together and when this happens it was found that they have changed somewhat their larval feeding habits. In the Boquillas Canyon, Dryden, and Langtry areas where gilberti and mariae fly together, gilberti larvae penetrate into the caudex of the plant to a depth of 20 mm, whereas mariae does not penetrate more than 5 mm into the caudex, but does most of its feeding in the lower portion of the leaves. In the Del Rio, Juno, and Bracketville areas where gilberti and rindgei fly together gilberti penetrates sometimes to a depth of 30 mm into the caudex, whereas rindgei seldom penetrates below 20 mm. It is not unusual to find these two species and estelleae (Stallings & Turner) feeding in the same plant, each with a somewhat different feeding level, with estelleae feeding only in the leaves, rindgei the base of the leaves and upper caudex, and gilberti in the lower base of the leaves and deeper into the caudex. A rea-

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sonable explanation for this is the rapidly declining population of *lecheguilla* plants in this general area, thus resulting in greater competition for food among these insects. The pH of the soil has considerable bearing on the occurrence of these species in a given habitat. *Mariae* is associated with distinctly alkaline soil, often from 7.7-8 pH; *chinatiensis* and *lajitaensis*, with soil somewhat less alkaline, around pH 7.2-7.4; and *gilberti* and *rindgei*, in soil that is neutral or only slightly alkaline, pH 7-7.1.

An interesting observation was made concerning the feeding habits of the larvae of Aegiale hesperiaris (Walker) and Agathymus hoffmanni (Freeman) in Agave americana L. in Mexico. In areas where *hoffmanni* is fairly abundant, especially in the valley of Mexico, the larvae feed in a small species of a parryilike Agave, and overlook the larger americana plants, in which the larvae of hesperiaris are found. The hoffmanni trap doors are located on the lower surface and usually near the center of the leaf. In areas where both species are found in the same plant, some member of the americana complex, hoffmanni prefer the hip of the leaf for their feeding area leaving the center portion for the much larger hesperiaris larvae, or else the hesperiaris forced them into that position. Specimens of hoffmanni from these areas are atypical, indicating that they have evolved or else are in the process of evolving into a separate species. Both biological and morphological work on these specimens is difficult due to the scarcity of the individuals. When sufficient material is present for study it may then be possible to determine the exact status of these hoffmanni-like specimens.

In Arizona some parallel evolution appears to be present in various populations of Agathymus aryxna (Dyar). In the general area of the Chiricahua Mountains southward through Ramsey Canyon or into northern Mexico is found one population with similar characteristics. Another population begins near Globe and extends through Santa Catalina, Rincon, and Santa Rita Mountains southward through the Patagonia Mountains into Sonora, Mexico. The third is a rather variable population that occurs in the Baboquivari Mountains and possibly on southward into Mexico. Indications are that at one time this entire area was covered with a palmeri-like Agave in which fed an aryxna-like prototype of our modern species. Environmental conditions plus man made factors eliminated the Agave in large areas thus producing desert and mountain islands where the plants are still undisturbed. Individuals from each population may be recognized by the width of the spots on both wings.

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Starting with the eastern population, the spots are generally separate and fairly small. In the central population, the spots are somewhat larger and tend to be closer together. While in the western population, the spots are the largest, tending to fuse together in some cases. To me there seems no doubt that we are seeing evolved three separate subspecies that eventually will assume specific status within a number of years.

In the neumoegeni complex we have a number of so called species that exhibit characteristics indicating a common ancestor. Possibly some of these have evolved farther than others and are specific, however others may still be in the process of changing from distinct subspecies into full species. The test of breeding has just started to be worked upon by Stallings, Turner & Stallings and the results will likely be very conclusive in forming our conception of this complex. I have collected larvae of wild hybrids of a cross between male mariae and female judithae (Stallings & Turner), as well as the same in a cross between male mariae and female diabloensis Freeman and the resulting F_1 did not resemble each other as much as the pure judithae and diabloensis do, especially the females. In all of the wild hybrids that I have seen to date involving mariae with some species of the neumoegeni complex it has always been the males of mariae that have resulted in the hybrids. This is based on the presence of the larvae in species of Agave other than members of the lecheguilla complex. From all evidence presented these F_1 are sterile.

Two members of the *neumoegeni* complex are associated with juvenile plants, *neumoegeni* (Edwards) and *florenceae* (Stallings & Turner). All other species are associated with mature plants running from small to large in size. *Florenceae* and *neumoegeni* morphologically resemble each other more closely than do any of the other species in this group even though there is a great distance separating them in their distribution. Genetically speaking the entire complex is closely related.

In the *remingtoni* complex we have three described species, *remingtoni* (Stallings & Turner), *estelleae* (Stallings & Turner) and *comstocki* (Harbison), and several undescribed species. Genetically they are related, biologically they are very different. *Remingtoni* is found in the mountains in very rugged terrain, often on cliffs, with the pH on the acid side, and feeding upon a member of the *lecheguilla* complex. *Estelleae* is associated with alkaline or near neutral soil, usually in the plains country or else

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where it is slightly hilly, and feeding upon another member of the *lecheguilla* complex. *Comstocki* is found in hilly, desert, areas of Baja California (pH has not been checked), feeding upon *Agave shawii* Engelm., which is not at all related to *lecheguilla*. The general range of *remingtoni* is Victoria, Tamps., Mexico southward to the Jacala, Hidalgo, area, while *estelleae* occurs north and west of Victoria. All indications point to a common ancestor of these two species that occurred in an area where the gene pool was available for all regions of the range. Due to environmental changes, both man made and natural, the gene pool was disrupted, eventually resulting in the formation of separate subspecies and eventually species.

In conclusion I would say that in the genus Agathymus we have had three major prototype complexes, the *neumoegeni*, *aryxna*, and *mariae*, and from these three all of our present day species evolved.

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